

# Tools and more for High-Performance Computing

## Overview

- ❑ Environment & account setup
- ❑ Compilers
- ❑ IDEs, Libraries
- ❑ Make & Makefiles
- ❑ Version control
- ❑ Data analysis tools: awk & perl (self study)
- ❑ Visualization tools (self study)
- ❑ Resource Managers

## The DTU computer system



January 2019

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3

## The central DTU UNIX system

- ❑ Application servers – x86\_64 based:
  - ❑ 8 Huawei XH620 V3 (2x Intel Xeon E5-2660v3 2.6 GHz)
  - ❑ 7 Dell PowerEdge FC430 (2x Intel Xeon E5-2670v3 2.3 GHz)
  - ❑ Scientific Linux 6.4
- ❑ Desktop servers (ThinLinc):
  - ❑ 4 servers (4x AMD Opteron 6376, 2.4 GHz)
- ❑ 10000+ users (students + employees)

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4

# The DTU computer system

## ❑ HPC servers:

- ❑ 64 HP SL2x170z (2x Xeon 5550 2.6 GHz, 24 GB memory)
- ❑ 42 IBM NeXtScale nx360 M4 (2x Xeon E5-2680v2 2.8 GHz, 128 GB memory)
- ❑ 40 Huawei XH620 V3 (2x Intel Xeon E5-2660v3 2.6 GHz)
- ❑ 38 Huawei XH620 V3 (2x Intel Xeon E5-2650v4 2.2 GHz) (**new setup – used in this course!**)

## ❑ + “private” clusters

- ❑ DTU Compute, DTU Nanotech, DTU Photonics, DTU Chemistry
- ❑ ...

# Access to the system

## ❑ Remote access, only:

### ❑ ThinLinc remote desktop session:

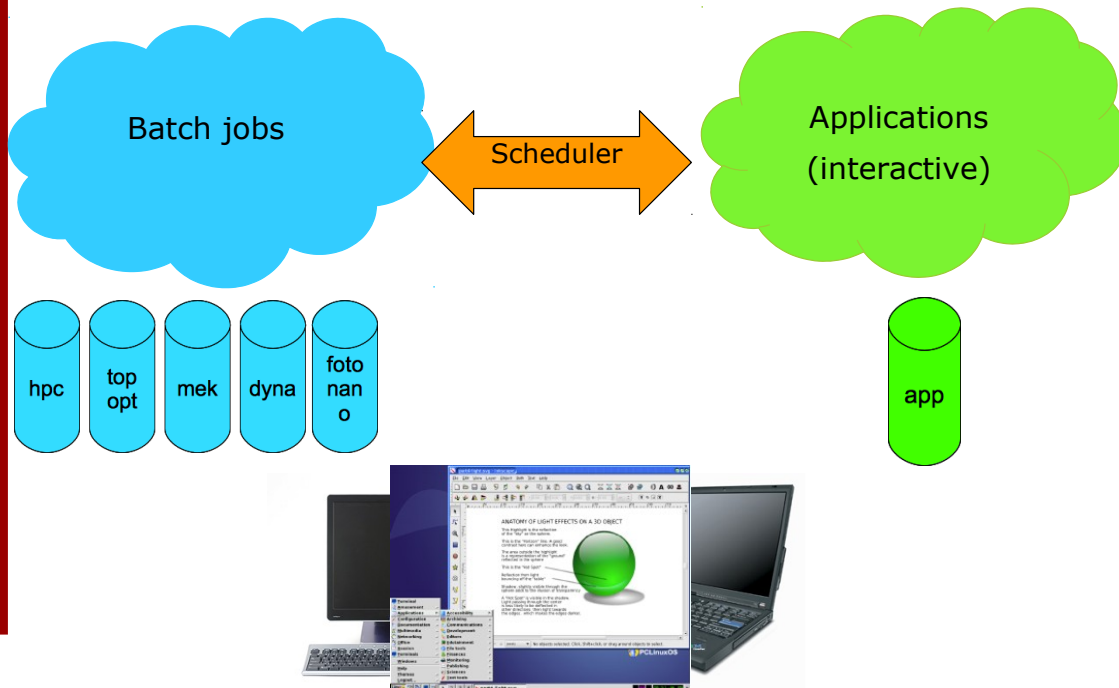
- ❑ download ThinLinc client from [www.thinlinc.com](http://www.thinlinc.com)
- ❑ connect to [thinlinc.gbar.dtu.dk](http://thinlinc.gbar.dtu.dk)
- ❑ preferred way, if you work a lot with GUIs
- ❑ browser based: <https://thinlinc.gbar.dtu.dk/>

### ❑ Secure SHell (ssh) connection ([login.hpc.dtu.dk](http://login.hpc.dtu.dk) or [login.gbar.dtu.dk](http://login.gbar.dtu.dk))

### ❑ On Campus:

- ❑ ThinLinc from Windows computers (Winbar).

## Our Setup



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7

## Access to Linux machines (02614)

- ❑ We have created a special environment for this course, only!
- ❑ Connect by SSH ([hpc2019.hpc.dtu.dk](https://hpc2019.hpc.dtu.dk))
  - ❑ linuxsh – access to interactive nodes
  - ❑ do not execute programs on the login node!
  - ❑ Note: X11 forwarding with '-X' option!
- ❑ or ThinLinc session ([hpc2019.gbar.dtu.dk](https://hpc2019.gbar.dtu.dk))
  - ❑ open a terminal (xterm-appnode) from the DTU menu – access to interactive node (same as linuxsh, above)

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8

## Access to Linux machines (02614)

The special setup consists of

- ❑ ThinLinc cluster (4 desktop nodes)
- ❑ 4 interactive nodes:
  - ❑ Lenovo SD530 server
  - ❑ 2x Intel Xeon Gold 6126 (2.6 GHz)
  - ❑ 192 GB memory
- ❑ 12 batch nodes in the LSF 'hpcintro' queue
  - ❑ same as above
  - ❑ 4 of the 12 nodes have 384GB memory

## The DTU computer system

Be aware of, that ...

- ❑ this is a multi-user system(!)
- ❑ (almost) all applications on the system are started by a load-balancing queueing system
- ❑ there are different
  - ❑ CPU types,
  - ❑ clock frequencies,
  - ❑ amounts of RAM,
  - ❑ etc

# The DTU computer system

Comparing performance numbers:

- ❑ make sure to be on the same machine type
  - ❑ `lscpu` command
  - ❑ `echo $CPULYPEV`
- ❑ check the load (interactive sessions)
  - ❑ `uptime` command
- ❑ check the # of CPU cores
  - ❑ `cpucount` command

## Compilers

- ❑ Oracle Studio compilers & tools
  - ❑ version 12 upd 6 ('module load studio')
  - ❑ commands: `suncc`, `sunCC`, `sunf95`
  - ❑ sets `cc`, `CC` and `f95`, `f77`, too!
- ❑ GNU Compilers (C/C++)
  - ❑ `gcc` 4.8.5 (OS standard)
  - ❑ `gcc` 6.3.0 ('module load gcc')
  - ❑ newer versions: check with 'module avail gcc'
- ❑ Note: 'cc' depends on the module loaded!!!
  - ❑ always use the specific names, i.e. `gcc`, `suncc`, ...

## More compilers

- ❑ Portland Group / PGI compilers
  - ❑ version 2013 ('module load pgi' – default)
  - ❑ version 2015 ('module load pgi/2015\_15.7')
  - ❑ version 2016 ('module load pgi/2016\_16.3')
  - ❑ version 2017 ('module load pgi/2017\_17.1')
  - ❑ version 2018 ('module load pgi/2018\_18.5')
  - ❑ commands: pgcc, pgCC, pgf95/pgf77

## More compilers

- ❑ Intel compilers
  - ❑ version 13.0.1 ('module load intel')
  - ❑ commands: icc, ifort
  - ❑ + some extra tools
  - ❑ other versions: check with 'module avail intel'
    - ❑ intel/2018.0.033
    - ❑ intel/2017.0.035
    - ❑ intel/2016.2.0
    - ❑ intel/2015.3.187
    - ❑ ...

## Using modules

- ❑ modules help to organize certain Unix environment settings, e.g. PATH, MANPATH, LD\_LIBRARY\_PATH, etc. for different versions of the same application
- ❑ list available modules: `module avail`
- ❑ load a module: `module load gurobi`
- ❑ swap a version: `module swap gurobi/5.6.3`
- ❑ swap to default: `module swap gurobi`
- ❑ info: <http://gbar.dtu.dk/index.php/faq/83-modules>

## IDEs

- ❑ Oracle Studio (`sunstudio`)
  - ❑ Compilers (Fortran, C/C++)
  - ❑ Debugger (`dbx`), analysis tools – more later
- ❑ Eclipse (`eclipse4`)
- ❑ Graphical debuggers:
  - ❑ Totalview (`totalview`)
  - ❑ Dbxtool (`dbxtool`)
  - ❑ Data Display Debugger (`ddd`)
    - ❑ GUI front-end to either `dbx` or `gdb`



# Libraries

- ❑ Available Scientific Libraries:
  - ❑ ATLAS
    - ❑ BLAS, CBLAS, LAPACK, ...
    - ❑ installed on all nodes
  - ❑ FFTW3
    - ❑ installed on all nodes
  - ❑ Solaris Studio Performance Library (optimized)
    - ❑ BLAS, LAPACK, FFT, ...
    - ❑ part of Oracle Studio
    - ❑ installed on all nodes

# Make & Makefiles

A tool for building and  
maintaining software projects

## Make – The ideas behind

- ❑ maintain, update and regenerate groups of programs
- ❑ useful tool in multi-source file software projects
- ❑ can be used for other tasks as well, e.g. typesetting projects, flat-file databases, etc
- ❑ in general: every task that involves updating files (i.e. result) from other files (i.e. sources) is a good candidate for make

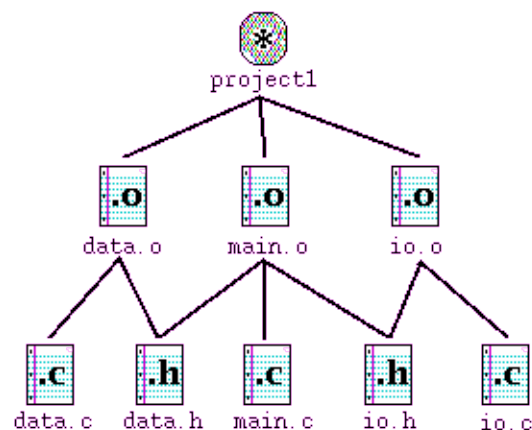
## Make – The ideas behind

Dependency graph:

result (executable)

intermediate level

source file level



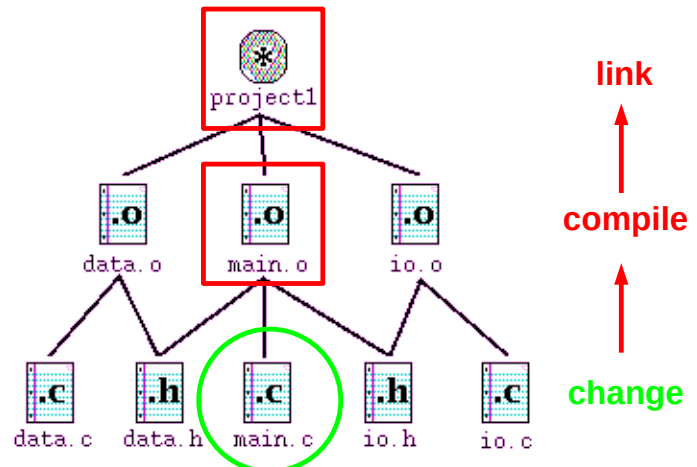
# Make – The ideas behind

Dependency graph:

result (executable)

intermediate level

source file level



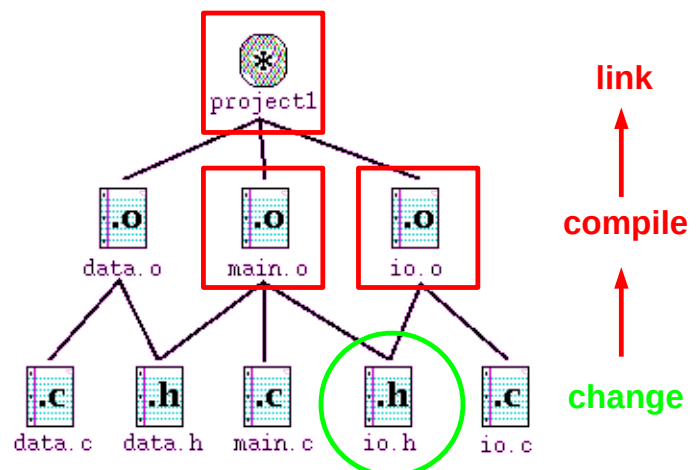
# Make – The ideas behind

Dependency graph:

result (executable)

intermediate level

source file level



## Make – The ideas behind

- ❑ Compiling by hand:
  - ❑ error prone
  - ❑ easy to forget a file
  - ❑ typos on the command line
- ❑ There is a tool that can help you:

**make**

## Make – The ideas behind

Things 'make' has to know:

- ❑ file status (timestamp)
- ❑ file location (source/target directories)
- ❑ file dependencies
- ❑ file generation rules (compiling/linking)
  - ❑ general rules ( `.c`  $\rightarrow$  `.o` )
  - ❑ special rules ( `io.c`  $\rightarrow$  `io.o` )
- ❑ tools (compilers, etc.)

- filesystem

- Makefile

- environment

## Makefile – rulesets...and more

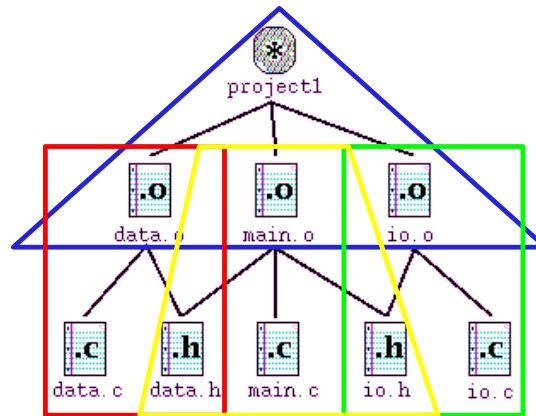
- ❑ make needs a set of rules to do its job
- ❑ rules are defined in a text file – the *Makefile*
- ❑ standard names: Makefile or makefile
- ❑ non-standard names can be used with the '-f' option of make: `make -f mymf ...`
- ❑ preview/dryrun option: `make -n ...`

## Makefile – rulesets...and more

There are two major object types in a Makefile

- ❑ **targets**
  - ❑ definition by a ":"
  - ❑ followed by the dependencies (same line)
  - ❑ followed by lines with the commands to execute
- ❑ **macros**
  - ❑ definition by "="
  - ❑ single line (use "\" to extend lines)
- ❑ ... and comments: (lines) starting with #

# Makefile – rulesets...and more



```
project1: data.o main.o io.o
    cc data.o main.o io.o -o project1
data.o: data.c data.h
    cc -c data.c
main.o: data.h io.h main.c
    cc -c main.c
io.o: io.h io.c
    cc -c io.c
```

# Makefile – rulesets...and more

target

dependencies

```
project1: data.o main.o io.o
    cc data.o main.o io.o \
    -o project1
    echo "Done."
```

command(s) to execute

TAB !!!

```
data.o: data.c data.h
    cc -c data.c
```

comment line

```
# the main program
main.o: data.h io.h main.c
    cc -c main.c
```

# Makefile – rulesets...and more

```
# Sample Makefile
CC      = gcc
OPT      = -g -O3
WARN     = -Wall
CFLAGS  = $(OPT) $(WARN)    # the C compiler flags
OBJECTS  = data.o main.o io.o
```

Macro definitions

```
project1 : $(OBJECTS)
    $(CC) $(CFLAGS) -o project1 $(OBJECTS)
```

Macro reference

```
clean:
    @rm -f *.o core
```

```
realclean : clean
    @rm -f project1
```

Where are my rules  
for compiling the .o files?

```
# file dependencies
data.o : data.c data.h
main.o : data.h io.h main.c
io.o   : io.h io.c
```

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29

# Makefile – rulesets...and more

## Running make:

```
myhost $ make
gcc -g -O3 -Wall -c -o data.o data.c
gcc -g -O3 -Wall -c -o main.o main.c
gcc -g -O3 -Wall -c -o io.o io.c
gcc -g -O3 -Wall -o project1 data.o main.o io.o
```

How did **make** know how to build data.o, ... ?

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30

# Makefile – rulesets...and more

built-in data base of “*standard rules*” and “*standard macros*”:

- ❑ known rules:
  - ❑ compile .o files from a .c/.cpp/.f/... source file
  - ❑ link executables from .o files
- ❑ pre-defined macros:
  - ❑ CC, CFLAGS, FC, FFLAGS, LD, LDFLAGS
- ❑ view with `make -p -f /dev/null`  
(long listing!)

# Makefile – rulesets...and more

```
# GNU Make 3.80
# Variables
...
# default
OUTPUT_OPTION = -o $@
# makefile (from `Makefile', line 3)
CC = gcc
# environment
MACHTYPE = i686-suse-linux
# makefile (from `Makefile', line 6)
CFLAGS = $(OPT) $(WARN)
# makefile (from `Makefile', line 4)
OPT = -g -O3
# makefile (from `Makefile', line 5)
WARN = -Wall
# default
COMPILE.c = $(CC) $(CFLAGS) $(CPPFLAGS) -c
# makefile (from `Makefile', line 8)
OBJECTS = data.o main.o io.o
...
```



## Makefile – rulesets...and more

```
...

# Implicit Rules
.c.o:
#  commands to execute (built-in):
#    $(COMPILE.c) $(OUTPUT_OPTION) $<

...

data.o: data.c data.c data.h
#  Implicit rule search has been done.
#  Implicit/static pattern stem: `data'
#  Last modified 2004-08-27 10:08:56.008831584
#  File has been updated.
#  Successfully updated.
#  commands to execute (built-in):
#    $(COMPILE.c) $(OUTPUT_OPTION) $<
```

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33

## Makefile – rulesets...and more

Practical hints:

- ❑ preview/dryrun option: `make -n ...`
- ❑ switch off built-in rules/macros:  
`make -r ...`
- ❑ check the known suffixes (.SUFFIXES) and implicit rules for your source files, e.g. does gmake still fail for .f90/.f95
- ❑ add suffixes needed:  
`.SUFFIXES: .f90`

January 2019

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34

## Makefile – rulesets...and more

Practical hints (cont'd):

- ❑ be aware of timestamps (Network-FS)
- ❑ override macros on the command line:

```
myhost $ make
gcc -g -O3 -Wall      -c -o data.o data.c
gcc -g -O3 -Wall      -c -o main.o main.c
gcc -g -O3 -Wall      -c -o io.o io.c
gcc -g -O3 -Wall-o project1 data.o main.o io.o
```

```
myhost $ make CFLAGS=-g
gcc -g      -c -o data.o data.c
gcc -g      -c -o main.o main.c
gcc -g      -c -o io.o io.c
gcc -g -o project1 data.o main.o io.o
```

## Makefile – rulesets...and more

Special variables/targets:

- ❑ the first target in Makefile is the one used when you call make without arguments!
- ❑ automatic variables:
  - ❑ \$< - The name of the first prerequisite.
  - ❑ \$@ - The file name of the target of the rule.
- ❑ for more information:
  - ❑ man make
  - ❑ info make

## Makefile – rulesets...and more

Makefile design – Best practice:

- ❑ start with the macros/variables
- ❑ call your first target “all:” and make it depend on all targets you want to build
- ❑ have a target “clean:” for cleaning up
- ❑ avoid explicit rules where possible, i.e. use redundancy

## Makefile – rulesets...and more

Makefile design – Best practice (cont'd):

- ❑ check your dependencies:
  - ❑ by hand
  - ❑ most C/C++ compilers can generate Makefile dependencies (see compiler documentation)
  - ❑ Sun Studio: `suncc -xM1`
  - ❑ Gnu C: `gcc -MM`
  - ❑ external tool: `makedepend -Y`
  - ❑ Note: the options above ignore `/usr/include`

## Makefile – rulesets...and more

Common mistakes:

- ❑ missing TAB in “command lines”
- ❑ wrong variable references:
  - \$VAR instead of \$(VAR)
- ❑ missing/wrong dependencies
- ❑ remember: each command is carried out in a new sub-shell

## Makefile – rulesets...and more

Makefiles – and Makefiles (from IDEs)

- ❑ Most IDEs create their own Makefiles
  - ❑ ... which are often not very smart
  - ❑ ... which are often not compatible
- ❑ make and (g)make:
  - ❑ Linux: make == gmake (GNU make)
  - ❑ Unix: make != gmake
  - ❑ if make fails, try gmake

## Make and Makefiles: Labs

- ❑ There are five short lab exercises
- ❑ download from Campusnet
- ❑ unzip the file
- ❑ the exercises are in the directories lab\_N
- ❑ read the README files for instructions

## Make and Makefiles: Labs

- ❑ Hints:
  - ❑ M\_PI is a definition from <math.h>
  - ❑ sin() is a function from libm.so, so you have to link with that library (use -lm the right place)

## Version control

- ❑ Larger – but also simple – software projects need to keep track of different versions
- ❑ This is very useful during development, e.g. to be able to go back to the last working version
- ❑ Versioning Tools:
  - ❑ RCS – single user, standalone
  - ❑ CVS – multi-user, network based
  - ❑ Subversion – multi-user, network based
  - ❑ git – multi-user, network based

## Version control

- ❑ DTU has a central CVS server
  - ❑ nice tool to share and control source files
  - ❑ request access on <https://repos.gbar.dtu.dk/>
  - ❑ basic introduction: <http://gbar.dtu.dk/faq/34-cvs>
- ❑ ... and a Subversion (SVN) server as well
  - ❑ request access on <https://repos.gbar.dtu.dk/>
  - ❑ basic introduction: <http://gbar.dtu.dk/faq/39-svn>
- ❑ ... and some info about Git and GitLab:
  - ❑ <http://gbar.dtu.dk/faq/41-git>
  - ❑ <http://www.gbar.dtu.dk/faq/94-gitlab>

## Data analysis tools

- ❑ Scientific software usually produces lots of data/datafiles
- ❑ There are good tools to do (a quick) analysis:
  - ❑ awk – standard UNIX/Linux tool
  - ❑ perl – (almost) standard on many platforms
- ❑ Both tools can be used
  - ❑ from the command line
  - ❑ with scripts

## Data analysis tools – awk

- ❑ awk operators:

Field reference:	\$
\$0: the whole line - \$n: the n-th field	
Increment or decrement:	++ --
Exponentiate:	^
Multiply, divide, modulus:	* / %
Add, subtract:	+ -
Concatenation:	(blank space)
Relational:	< <= > >= != ==
Match regular expression:	~ !~
Logical:	&&
C-style assignment:	= += -= *= /= %= ^=

## Data analysis tools – awk

### Examples:

- ❑ Print first two fields in opposite order:

```
awk '{ print $2, $1 }' file
```

- ❑ Print column 3 if column 1 > column 2:

```
awk '$1 > $2 {print $3}' file
```

- ❑ Print line (default action) if col. 3 > col. 2:

```
awk '$3 > $2' file
```

## Data analysis tools – awk

### Examples (cont'd):

- ❑ Add up first column, print sum and average:

```
awk '{s += $1}; END { print "sum  
is", s, " avg is", s/NR}' file
```

- ❑ Special keywords/variables:

BEGIN	do before the first record
END	do after the last record
NR	number of records
NF	number of fields
\$NF	the value of the last field



## Data analysis tools

- ❑ Other useful standard Unix tools for data analysis:
  - ❑ sort
  - ❑ uniq
  - ❑ head, tail
  - ❑ wc
  - ❑ sed
  - ❑ ...

## Data analysis tools – perl

- ❑ Perl is a very powerful tool, that combines the features of awk, grep, sed, sort, and other Unix-tools into one language
- ❑ Good tool for more complex data analysis tasks
- ❑ Web-site: <http://perl.org/>
- ❑ Archive of perl programs:
  - ❑ Comprehensive Perl Archive Network – CPAN
  - ❑ <http://www.cpan.org/>

# Data analysis tools – perl

## Perl example script:

```
#!/usr/bin/perl

while (<>) {

    next if /^#/;          # skip comment lines
    @fields = split();     # split the line

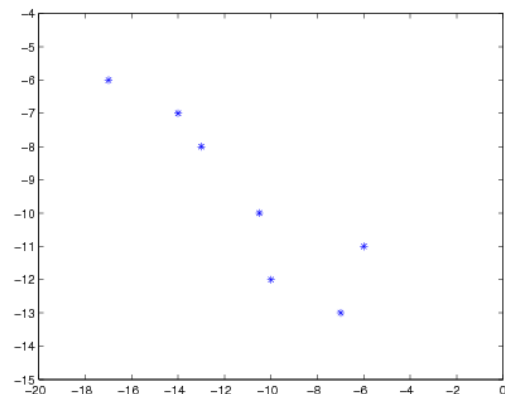
    if ($#fields == 2) {   # 3(!) elements
        print "$fields[0] $fields[2]\n";
    }
    else {
        print;
    }
}
```

# Visualization

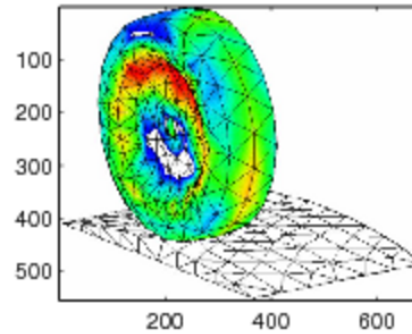
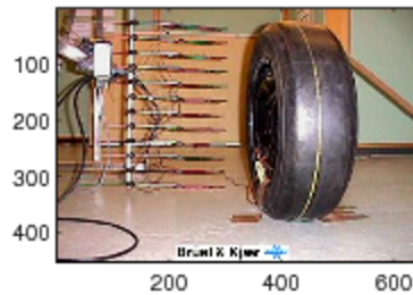
Visualization is an important part of Scientific Computing

Motivation: What's that?

```
A ( -17, -6)
B ( -14, -7)
C ( -13, -8)
D (-10.5, -10)
E (  -6, -11)
F (  -7, -13)
G ( -10, -12)
```



# Visualization



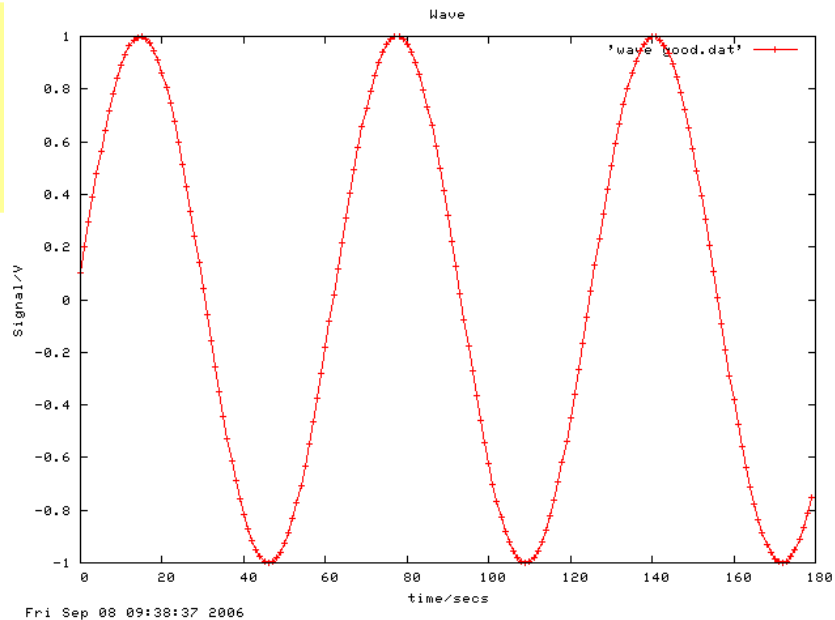
# Visualization

- ❑ Simple tools to visualize data:
  - ❑ Gnuplot (`gnuplot`)
    - ❑ command based, flexible
    - ❑ good for scripting, batch analysis
    - ❑ limited graphics (not always suitable for publishing)
  - ❑ Grace (`xmgrace`)
    - ❑ GUI-based
    - ❑ difficult to do scripting, batch analysis
    - ❑ very good graphics (publication-ready)
  - ❑ ... or whatever tool you like/prefer

# Visualization

## Gnuplot example:

```
gnuplot>  
gnuplot>  
gnuplot>  
gnuplot>  
gnuplot>
```



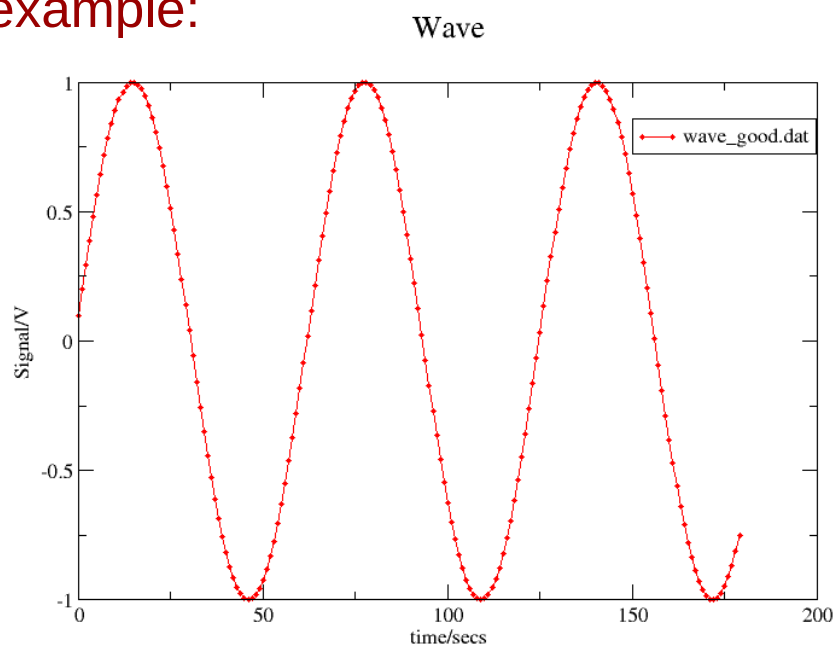
January 2019

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55

# Visualization

## Grace example:



Fri Sep 8 09:47:12 2006

January 2019

02614 - High-Performance Computing

56

# Visualization

- ❑ Best practice:
  - ❑ label the axes
  - ❑ use legends (and titles)
  - ❑ use the right scaling
    - ❑ a plot of a circle should be a circle
  - ❑ don't overload figures with information – use more figures instead
  - ❑ colors are useful – but can also be confusing

## And not to forget ...

... a very powerful tool/language for Scientific Computing:

- ❑ Python
  - ❑ built-in vector and matrix types (NumPy, SciPy)
  - ❑ data plot functionality (matplotlib)
  - ❑ interfaces to different languages
  - ❑ GPU support (PyOpenCL, PyCUDA)
  - ❑ and, and, and ....

## Data analysis – lab exercise

- ❑ download the file wave.zip from Campusnet
- ❑ follow the instructions in wave.readme
- ❑ Goal:
  - ❑ get used to awk (choose perl, if you like or know it already)
  - ❑ get used to either Gnuplot or Grace (or the tool you know/like)

## Resource Managers

To handle the workload on an HPC installation, one needs a tool to manage and assign the resources: a Resource Manager – sometimes also called 'batch queue system'

- ❑ Most common systems:
  - ❑ Torque/PBS (ext. scheduler, like Maui or MOAB)
  - ❑ LSF
  - ❑ Grid Engine
  - ❑ Slurm

# Resource Managers

Before submitting a job, one has to specify the resources needed, e.g.

- ❑ # of CPUs/cores
- ❑ amount of memory
- ❑ expected run time (wall-clock time)
- ❑ other resources, like disk space, GPUs, etc

This is done in a special job script and is system (RM) dependent – but very similar for all RMs.

# Resource Managers

The simplest job script:

```
#!/bin/bash
sleep 60
```

submit.sh

```
$ bsub < submit.sh
```

```
Job <702572> is submitted to default queue <hpc>.
```

```
$ bstat
```

JOBID	USER	QUEUE	JOB_NAME	SLOTS	STAT	START_TIME	ELAPSED
702572	gbarbd	hpc	NONAME	1	RUN	Dec 13 12:17	0:00:00

```
$ bjobs
```

JOBID	USER	QUEUE	JOB_NAME	SLOTS	STAT	START_TIME	TIME_LEFT
702572	gbarbd	hpc	NONAME	1	RUN	Dec 13 12:17	00:15:00 L

```
$ ls -g
```

```
total 4
```

-rw-r--r--	1	gbar	1493	Dec 13 12:18	NONAME_702572.out
-rw-r--r--	1	gbar	22	Dec 13 12:05	simple.sh

# Resource Managers

## The simplest job script – the full story:

```
#!/bin/bash
sleep 60
```

simple.sh

```
$ bsub < simple.sh
bsub info: Job has no name! Setting it to NONAME!
bsub info: Job has no wall-clock time! Setting it to 15 minutes!
bsub info: Job has no output file! Setting it to NONAME_%J.out!
bsub info: Job has no memory requirements! Setting it to 1024 MB!
bsub info: You need to specify at least -R "rusage[mem=...]"!
Job <702608> is submitted to default queue <hpc>.
```

# Resource Managers

## A simple job script:

```
#!/bin/bash
#BSUB -J sleeper
#BSUB -o sleeper_%J.out
#BSUB -q hpcintro
#BSUB -W 2
#BSUB -R "rusage[mem=512MB]"
```

```
sleep 60
```

```
$ bsub < submit.sh
Job <702645> is submitted to queue <hpcintro>.
```

```
$ ls -g
```

```
total 3
```

```
-rw-r--r-- 1 gbar 121 Dec 13 12:32 submit.sh
```

```
-rw-r--r-- 1 gbar 1592 Dec 13 12:36 sleeper_702646.out
```



# Resource Managers

## ❑ The output file:

```
Sender: LSF System <lsfadmin@n-62-21-20>
Subject: Job 702646: <sleeper> in cluster <dcc> Done

Job <sleeper> was submitted from host <hpclogin3> by user <gbarbd> in
cluster <dcc> at Wed Dec 13 12:34:59 2017.
Job was executed on host(s) <n-62-21-20>, in queue <hpc>, as user
<gbarbd> in cluster <dcc> at Wed Dec 13 12:34:59 2017.
</zhome/../../../../> was used as the home directory.
</zhome/../../../../02614/Batch/LSF> was used as the working directory.
Started at Wed Dec 13 12:34:59 2017.
Terminated at Wed Dec 13 12:36:00 2017.
Results reported at Wed Dec 13 12:36:00 2017.
```

Your job looked like:

```
-----
# LSBATCH: User input
#!/bin/bash
#BSUB -J sleeper
#BSUB -o sleeper_%J.out
```

# Resource Managers

## ❑ The output file (cont'd):

### ❑ job summary

Successfully completed.

Resource usage summary:

CPU time :	0.28 sec.
Max Memory :	4 MB
Average Memory :	4.00 MB
Total Requested Memory :	512.00 MB
Delta Memory :	508.00 MB
Max Swap :	-
Max Processes :	4
Max Threads :	5
Run time :	65 sec.
Turnaround time :	61 sec.

The output (if any) is above this job summary.

# Resource Managers

## Separating output and errors:

```
#!/bin/bash
#BSUB -J sleeper
#BSUB -o sleeper_%J.out
#BSUB -e sleeper_%J.err
#BSUB -q hpcintro
#BSUB -W 2 -R "rusage[mem=512MB]"

rm nonexistent.txt
echo "Just a minute ..."
sleep 60

$ bsub < submit2.sh

...
$ ls -g
total 3
-rw-r--r-- 1 gbar 184 Dec 13 13:56 submit2.sh
-rw-r--r-- 1 gbar 63 Dec 13 13:59 sleeper_702793.err
-rw-r--r-- 1 gbar 1744 Dec 13 14:00 sleeper_702793.out
```

# Resource Managers

## Separating output, errors – and mail summary:

```
#!/bin/bash
#BSUB -J sleeper
#BSUB -o sleeper_%J.out
#BSUB -e sleeper_%J.err
#BSUB -q hpcintro
#BSUB -W 2 -R "rusage[mem=512MB]"

rm nonexistent.txt
echo "Just a minute ..."
sleep 60

$ bsub -N < submit2.sh

...
$ ls -g
total 3
-rw-r--r-- 1 gbar 184 Dec 13 13:56 submit2.sh
-rw-r--r-- 1 gbar 63 Dec 13 14:04 sleeper_702814.err
-rw-r--r-- 1 gbar 18 Dec 13 14:04 sleeper_702814.out
```

send summary  
at end of job

# Resource Managers

A simple parallel job script:

- ❑ for OpenMP (single node), using 4 cores

```
#!/bin/bash
#BSUB -J openmp_para
#BSUB -o openmp_para_%J.out
#BSUB -q hpcintro
#BSUB -W 2 -R "rusage[mem=512MB] "
#BSUB -n 4 -R "span[hosts=1] "

export OMP_NUM_THREADS=$LSB_DJOB_NUMPROC
...
```

# Resource Managers

Another parallel job script:

- ❑ for MPI: two nodes, using 4 cores/node

```
#!/bin/bash
#BSUB -J mpi_para
#BSUB -o mpi_para_%J.out
#BSUB -q hpcintro
#BSUB -W 2 -R "rusage[mem=512MB] "
#BSUB -n 8 -R "span[ptile=4] "

module load mpi
mpirun ...
```

## Resource Managers

more options and examples:

- ❑ see <http://www.hpc.dtu.dk/> under
  - ❑ LSF User Guides
    - ❑ [http://www.hpc.dtu.dk/?page\\_id=2534](http://www.hpc.dtu.dk/?page_id=2534)
- ❑ do the lab exercises
- ❑ use 'man bsub', 'man bjobs', etc

## Resource Managers

DTU Computing Center specific commands:

- ❑ bstat – shows the status of your jobs; use 'bstat -h' for help for other options
- ❑ classstat – shows the status of the queues, e.g. free and used cores, pending jobs, etc
- ❑ nodestat – shows the current status of all nodes (use 'nodestat hpc' for the nodes of the 'hpc' queue)
- ❑ all commands above have a help (-h) option, but no man-page!

# Resource Managers

- ❑ There are a few hands-on exercises on CampusNet to get you acquainted with the batch system
- ❑ more information can be found on [www.hpc.dtu.dk](http://www.hpc.dtu.dk) under **LSF User Guides**
- ❑ we have a special queue for this course, 'hpcintro', so please use '-q hpcintro' instead of '-q hpc' in your job scripts!